IN THE SPECIFICATION

Page 3, replace the paragraph starting on line 21 with the following:

Figure 1: Partial sequence alignment of the -1 binding pocket motif of *Sulfolobus solfataricus* β-glycosidase (SSβG) (Cubellis *et al.*, *supra*) with high sequence similarity (left hand column gives SWISSPROT or TrEMBL annotation, numbering is that of SSβG); glycosidases with similar substrate specificity (a) to SSβG and glycosidases with different and/or broadened specificities in which E432 (d), W433 (c) and M439 (b, c, d) differ (marked with arrow and highlighted) (*Dalbergia cochinchinensis* β-glucosidase - Cairns *et al.,* TREMBL Accession No. Q9SPK3; *Costus speciosus* furostanol-β-glycoside hydrolase - Inoue *et al., FEBS Lett.* **389** (1996) 273-277; LPH_HUMAN, human lactase phlorizin hydrolase - Mantei *et al., EMBO J.,* **7** (1988) 2705-2713; MY3_SINAL, myrosinase from *Sinapsis alba* - Xue *et al., Plant Mol. Biol.,* **18** (1992) 387-398; LACG_STAAU (6-PBG), *S. aureus* 6-phosphogalactosidase - Breidt and Stewart, *Appl. Environ. Microbiol.,* **53** (1987) 969-973). The amino acid sequences are presented as amino acids 425 to 450 of SEQ ID NO: 2 and SEQ ID NOS: 3 to 18.